0550 0405 OIPE

**CRF Errors Corrected by the STIC Systems Branch**

**Serial Number:** 09/901,556A ENTERED

CRF Processing Date: 4/9/2002
Edited by:
Verified by: _____ (STIC staff)

- ☐ Changed a file from non-ASCII to ASCII
- ☑ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically:

  _____
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____.
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically:

  _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:

  _____
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:

  _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included:

  _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically:

  _____
- ☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____.
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically:

  _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically:

  _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted ***ending*** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☐ Other:

  _____

  _____

  _____

  _____

***Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.**

3/1/95

RAW SEQUENCE LISTING DATE: 04/09/2002
PATENT APPLICATION: US/09/901,556A TIME: 17:34:37

Input Set : A:\PTO.AMC.txt
Output Set: N:\CRF3\04092002\I901556A.raw

<110> APPLICANT: Hotten, Gertrud
Neidhardt, Helge
Bechtold, Rolf
Pohl, Jens
<120> TITLE OF INVENTION: GROWTH/DIFFERENTIATION FACTORS OF THE TGF-B FAMILY
<130> FILE REFERENCE: 100564-09021
<140> CURRENT APPLICATION NUMBER: 09/901,556A
<141> CURRENT FILING DATE: 1999-08-25
<150> PRIOR APPLICATION NUMBER: 08/289,222
<151> PRIOR FILING DATE: 1994-08-12
<150> PRIOR APPLICATION NUMBER: DE P 44 23 190.3
<151> PRIOR FILING DATE: 1994-07-01
<150> PRIOR APPLICATION NUMBER: EPO 92102324.8
<151> PRIOR FILING DATE: 1992-02-12
<150> PRIOR APPLICATION NUMBER: PCT/EP93/00350
<151> PRIOR FILING DATE: 1993-02-12
<160> NUMBER OF SEQ ID NOS: 53
<170> SOFTWARE: PatentIn version 3.1
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<213> ORGANISM: Homo sapiens
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gccgtttcgc ccacccccca tcacacccca cgagtacatg ctctcgctgt acaggacgct 240
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82 <212> TYPE: DNA
83 <213> ORGANISM: Homo sapiens
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90 cccagcaatg acctcctcat tgcttctggc ctttctcctc ctggctccaa ccacagtggc 180
92 cactcccaga gctggcggtc agtgtccagc atgtgggggg cccaccttgg aactggagag 240
94 ccagcgggag ctgcttcttg atctggccaa gagaagcatc ttggacaagc tgcacctcac 300
96 ccagcgccca acactgaacc gccctgtgtc cagagctgct ttgaggactg cactgcagca 360
98 cctccacggg gtcccacagg gggcacttct agaggacaac agggaacagg aatgtgaaat 420
100 catcagcttt gctgagacag gcctctccac catcaaccag actcgtcttg attttcactt 480
102 ctcctctgat agaactgctg gtgacaggga ggtccagcag gccagtctca tgttctttgt 540
104 gcagctccct tccaatacca cttggacctt gaaagtgaga gtccttgtgc tgggtccaca 600
106 taataccaac ctcaccttgg ctactcagta cctgctggag gtggatgcca gtggctggca 660
108 tcaactcccc ctagggcctg aagctcaagc tgcctgcagc caggggcacc tgaccctgga 720
110 gctggtactt gaaggccagg tagcccagag ctcagtcatc ctgggtggag ctgcccatag 780
112 gccttttgtg gcagcccggg tgagagttgg gggcaaacac cagattcacc gacgaggcat 840
114 cgactgccaa ggagggtcca ggatgtgctg tcgacaagag ttttttgtgg acttccgtga 900
116 gattggctgg cacgactgga tcatccagcc tgagggctac gccatgaact tctgcatagg 960
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122 acccacggcc cggcgccccc tgtctctgct ctattatgac agggacagca acattgtcaa 1140
124 gactgacata cctgacatgg tagtagaggc ctgtgggtgc agttagtcta tgtgtggtat 1200
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128 gggaatgacc tcattctctg tccagaatgt ggactccctc ttcctgagca tcttatggaa 1320
130 attaccccac ctttgacttg aagaaacctt catctaaagc aagtcactgt gccatcttcc 1380
132 tgaccactac cctctttcct agggcatagt ccatcccgct agtccatccc gctagcccca 1440
134 ctccagggac tcagacccat ctccaaccat gagcaatgcc atctggttcc caggcaaaga 1500
136 caccettagc tcacctttaa tagaccccat aacccactat gccttcctgt cctttctact 1560
138 caatggtccc cactccaaga tgagttgaca caaccccttc ccccaatttt tgtggatctc 1620
140 cagagaggcc cttctttgga ttcaccaaag tttagatcac tgctgcccaa aatagaggct 1680
142 tacctacccc cctctttgtt gtgagccccc gtccttctta gttgtccagg tgaactacta 1740
144 aagctctctt tgcatacctt catccatttt ttgtccttct ctgcctttct ctatgccctt 1800
146 aaggggtgac ttgcctgagc tctatcacct gagctcccct gccctctggc ttcctgctga 1860
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150 gtggctattc tgtgtcccta cactacctgg ctacccccct ccatggcccc agctctgcct 1980
152 acattctgat ttttttttt ttttttttt tgaaaagtta aaaattcctt aatttttat 2040
154 tcctggtacc actaccacaa tttacagggc aatatacctg atgtaatgaa aagaaaaaga 2100
156 aaaagacaaa gctacaacag ataaaagacc tcaggaatgt acatctaatt gacactacat 2160
158 tgcattaatc aatagctgca ctttttgcaa actgtggcta tgacagtcct gaacaagaag 2220
160 ggtttcctgt ttaagctgca gtaacttttc tgactatgga tcatcgttcc tt 2272
163 <210> SEQ ID NO: 3
164 <211> LENGTH: 401
165 <212> TYPE: PRT
166 <213> ORGANISM: Homo sapiens
168 <400> SEQUENCE: 3

RAW SEQUENCE LISTING DATE: 04/09/2002
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Input Set : A:\PTO.AMC.txt
Output Set: N:\CRF3\04092002\I901556A.raw

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171 1               5                   10                  15
174 Ala Thr Ala Arg Thr Val Thr Pro Lys Gly Gln Leu Pro Gly Gly Lys
175             20                  25                  30
178 Ala Pro Pro Lys Ala Gly Ser Val Pro Ser Ser Phe Leu Leu Lys Lys
179         35                  40                  45
182 Ala Arg Glu Pro Gly Pro Pro Arg Glu Pro Lys Glu Pro Phe Arg Pro
183     50                  55                  60
186 Pro Pro Ile Thr Pro His Glu Tyr Met Leu Ser Leu Tyr Arg Thr Leu
187 65                  70                  75                  80
190 Ser Asp Ala Asp Arg Lys Gly Gly Asn Ser Ser Val Lys Leu Glu Ala
191                 85                  90                  95
194 Gly Leu Ala Asn Thr Ile Thr Ser Phe Ile Asp Lys Gly Gln Asp Asp
195             100                 105                 110
198 Arg Gly Pro Val Val Arg Lys Gln Arg Tyr Val Phe Asp Ile Ser Ala
199         115                 120                 125
202 Leu Glu Lys Asp Gly Leu Leu Gly Ala Glu Leu Arg Ile Leu Arg Lys
203     130                 135                 140
206 Lys Pro Ser Asp Thr Ala Lys Pro Ala Ala Pro Gly Gly Gly Arg Ala
207 145                 150                 155                 160
210 Ala Gln Leu Lys Leu Ser Ser Cys Pro Ser Gly Arg Gln Pro Ala Ser
211                 165                 170                 175
214 Leu Leu Asp Val Arg Ser Val Pro Gly Leu Asp Gly Ser Gly Trp Glu
215             180                 185                 190
218 Val Phe Asp Ile Trp Lys Leu Phe Arg Asn Phe Lys Asn Ser Ala Gln
219         195                 200                 205
222 Leu Cys Leu Glu Leu Glu Ala Trp Glu Arg Gly Arg Ala Val Asp Leu
223     210                 215                 220
226 Arg Gly Leu Gly Phe Asp Arg Ala Ala Arg Gln Val His Glu Lys Ala
227 225                 230                 235                 240
230 Leu Phe Leu Val Phe Gly Arg Thr Lys Lys Arg Asp Leu Phe Phe Asn
231                 245                 250                 255
234 Glu Ile Lys Ala Arg Ser Gly Gln Asp Asp Lys Thr Val Tyr Glu Tyr
235             260                 265                 270
238 Leu Phe Ser Gln Arg Arg Lys Arg Arg Ala Pro Leu Ala Thr Arg Gln
239         275                 280                 285
242 Gly Lys Arg Pro Ser Lys Asn Leu Lys Ala Arg Cys Ser Arg Lys Ala
243     290                 295                 300
246 Leu His Val Asn Phe Lys Asp Met Gly Trp Asp Asp Trp Ile Ile Ala
247 305                 310                 315                 320
250 Pro Leu Glu Tyr Glu Ala Phe His Cys Glu Gly Leu Cys Glu Phe Pro
251                 325                 330                 335
254 Leu Arg Ser His Leu Glu Pro Thr Asn His Ala Val Ile Gln Thr Leu
255             340                 345                 350
258 Met Asn Ser Met Asp Pro Glu Ser Thr Pro Pro Thr Cys Cys Val Pro
259         355                 360                 365
262 Thr Arg Leu Ser Pro Ile Ser Ile Leu Phe Ile Asp Ser Ala Asn Asn
263     370                 375                 380
266 Val Val Tyr Lys Gln Tyr Glu Asp Met Val Val Glu Ser Cys Gly Cys
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PATENT APPLICATION: US/09/901,556A TIME: 17:34:37

Input Set : A:\PTO.AMC.txt
Output Set: N:\CRF3\04092002\I901556A.raw

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274 <210> SEQ ID NO: 4
275 <211> LENGTH: 352
276 <212> TYPE: PRT
277 <213> ORGANISM: Homo sapiens
279 <400> SEQUENCE: 4
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286             20                  25                  30
289 Thr Leu Glu Leu Glu Ser Gln Arg Glu Leu Leu Leu Asp Leu Ala Lys
290         35                  40                  45
293 Arg Ser Ile Leu Asp Lys Leu His Leu Thr Gln Arg Pro Thr Leu Asn
294     50                  55                  60
297 Arg Pro Val Ser Arg Ala Ala Leu Arg Thr Ala Leu Gln His Leu His
298 65                  70                  75                  80
301 Gly Val Pro Gln Gly Ala Leu Leu Glu Asp Asn Arg Glu Gln Glu Cys
302                 85                  90                  95
305 Glu Ile Ile Ser Phe Ala Glu Thr Gly Leu Ser Thr Ile Asn Gln Thr
306             100                 105                 110
309 Arg Leu Asp Phe His Phe Ser Ser Asp Arg Thr Ala Gly Asp Arg Glu
310         115                 120                 125
313 Val Gln Gln Ala Ser Leu Met Phe Phe Val Gln Leu Pro Ser Asn Thr
314     130                 135                 140
317 Thr Trp Thr Leu Lys Val Arg Val Leu Val Leu Gly Pro His Asn Thr
318 145                 150                 155                 160
321 Asn Leu Thr Leu Ala Thr Gln Tyr Leu Leu Glu Val Asp Ala Ser Gly
322                 165                 170                 175
325 Trp His Gln Leu Pro Leu Gly Pro Glu Ala Gln Ala Ala Cys Ser Gln
326             180                 185                 190
329 Gly His Leu Thr Leu Glu Leu Val Leu Glu Gly Gln Val Ala Gln Ser
330         195                 200                 205
333 Ser Val Ile Leu Gly Gly Ala Ala His Arg Pro Phe Val Ala Ala Arg
334     210                 215                 220
337 Val Arg Val Gly Gly Lys His Gln Ile His Arg Arg Gly Ile Asp Cys
338 225                 230                 235                 240
341 Gln Gly Gly Ser Arg Met Cys Cys Arg Gln Glu Phe Phe Val Asp Phe
342                 245                 250                 255
345 Arg Glu Ile Gly Trp His Asp Trp Ile Ile Gln Pro Glu Gly Tyr Ala
346             260                 265                 270
349 Met Asn Phe Cys Ile Gly Gln Cys Pro Leu His Ile Ala Gly Met Pro
350         275                 280                 285
353 Gly Ile Ala Ala Ser Phe His Thr Ala Val Leu Asn Leu Leu Lys Ala
354     290                 295                 300
357 Asn Thr Ala Ala Gly Thr Thr Gly Gly Gly Ser Cys Cys Val Pro Thr
358 305                 310                 315                 320
361 Ala Arg Arg Pro Leu Ser Leu Leu Tyr Tyr Asp Arg Asp Ser Asn Ile
362                 325                 330                 335
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RAW SEQUENCE LISTING DATE: 04/09/2002
PATENT APPLICATION: US/09/901,556A TIME: 17:34:37

Input Set : A:\PTO.AMC.txt
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370 <211> LENGTH: 265
371 <212> TYPE: DNA
372 <213> ORGANISM: Homo sapiens
374 <400> SEQUENCE: 5
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379 cacagctgca ggcaccactg gaggggctc atgctgtgta cccacggccc ggcgcccct 180
381 gtctctgctc tattatgaca gggacagcaa cattgtcaag actgacatac ctgacatggt 240
383 agtagaggcc tgtgggtgca gttag 265
386 <210> SEQ ID NO: 6
387 <211> LENGTH: 139
388 <212> TYPE: DNA
389 <213> ORGANISM: Homo sapiens
391 <400> SEQUENCE: 6
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396 cgagtccaca ccacccacc 139
399 <210> SEQ ID NO: 7
400 <211> LENGTH: 27
401 <212> TYPE: DNA
402 <213> ORGANISM: Homo sapiens
404 <400> SEQUENCE: 7
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409 <211> LENGTH: 30
410 <212> TYPE: DNA
411 <213> ORGANISM: Homo sapiens
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417 <210> SEQ ID NO: 9
418 <211> LENGTH: 9
419 <212> TYPE: PRT
420 <213> ORGANISM: Homo sapiens
422 <400> SEQUENCE: 9
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425 1 5
428 <210> SEQ ID NO: 10
429 <211> LENGTH: 10
430 <212> TYPE: PRT
431 <213> ORGANISM: Homo sapiens
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436 1 5 10
439 <210> SEQ ID NO: 11
440 <211> LENGTH: 44
441 <212> TYPE: DNA